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KIG. 1C. HIG. 1d. **~** William. Selection 0 Steak. PARTY. 9 ω KIG. 10. i, e White Queen 5 3 Sand Second C) die. 88 SEA.

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_FIG. 2.

PURIFIED EXTRACELLULAR. PROTEINS STUDIED APPARENT MW 8Y SDS-PAGE (KD) S AMINO ACIDS S AMINO ACIDS S AMINO ACIDS NSKSV RO TORVS *71 ARAVG SR TEKTP AS DPEPA *32A FSRPG 32B FSRPG 32B FSRPG 24 APYEN 23.5 APKTY *16 AYPIT 14 ADPRL 12 FDTRL		
BY SDS-PAGE (KD) 5 AMINO ACIDS JSO NSKSV BO TDRVS *71 ARAVG 58 TEKTP A5 DPEPA *32A FSRPG 32B FSRPG 32B FSRPG 24 APYEN 23.5 APKTY *16 AYPIT JA ADPRL		
80 TDRVS *71 ARAVG 58 TEKTP 45 DPEPA *32A F5RPG 32B F5RPG *30 F5RPG 24 APYEN 23.5 APKTY *23 AETYL *16 AYPIT 14 ADPRL		
	80 *71 58 45 *32A 32B *30 24 23.5 *23 *16 14	TDRVS ARAVG TEKTP DPEPA FSRPG FSRPG APYEN APKTY AETYL AYPIT ADPRL

_FIG. 3.

EXTENDED N-TERMINAL SEQUENCE OF 30/32 KD COMPLEX OF M. TUBERCULOSIS EXTRACELLULAR PROTEINS

	10 20
30	FSRPGLPVEYLQVPSPSMGR
32 A	
32B	
1	H
1	21 30 40
30	DIKVQFQSGGNNSPAVYLLD
32 A	
32B	
	·

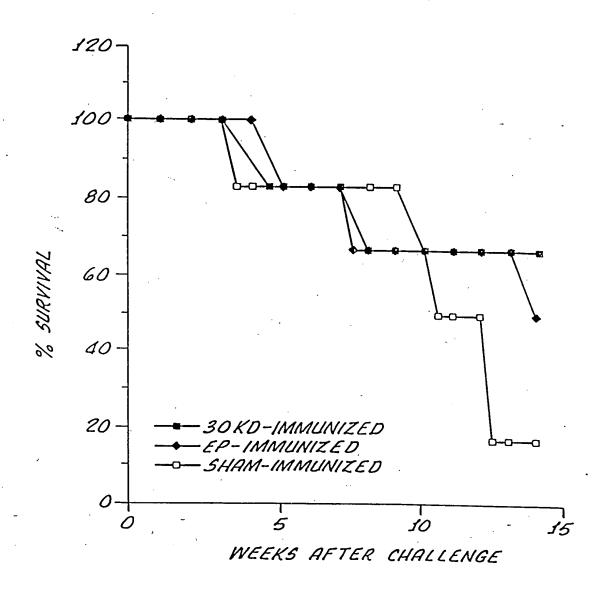
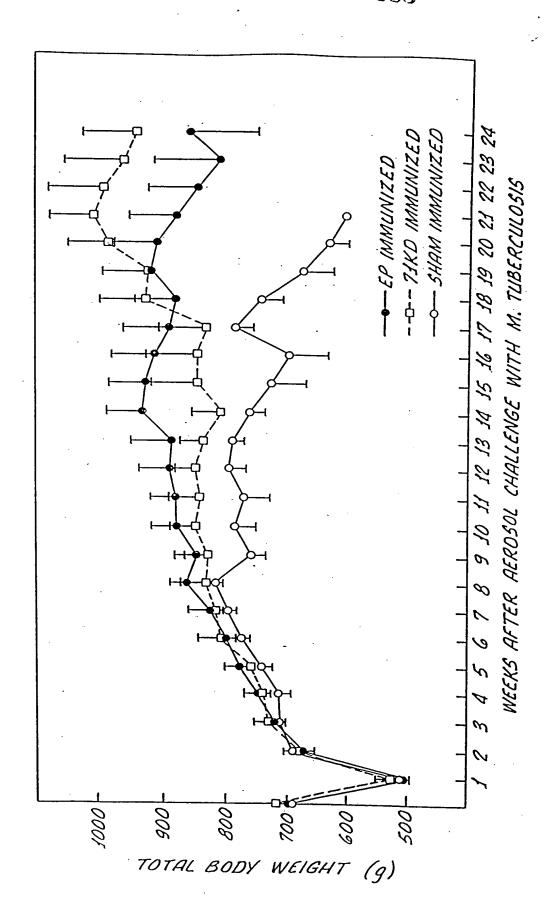
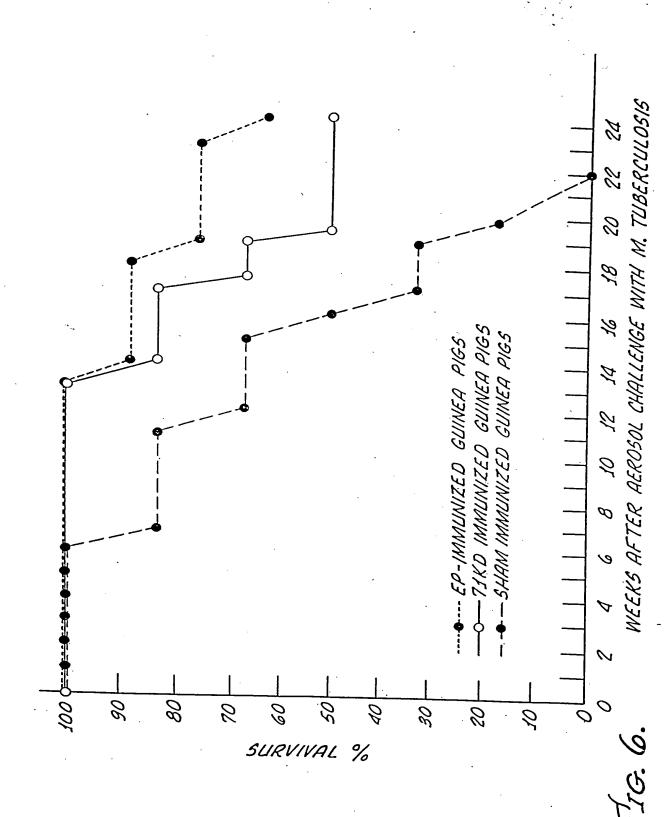


FIG. 4.



KTG. 5.



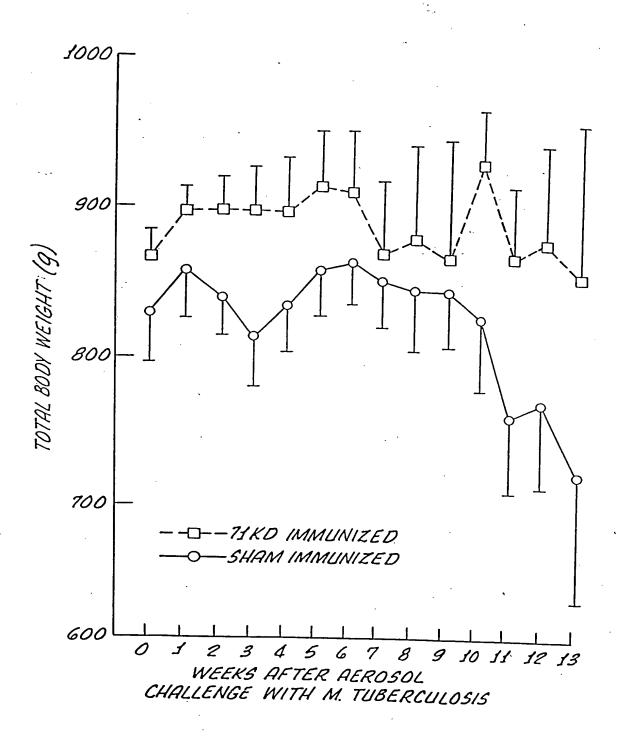
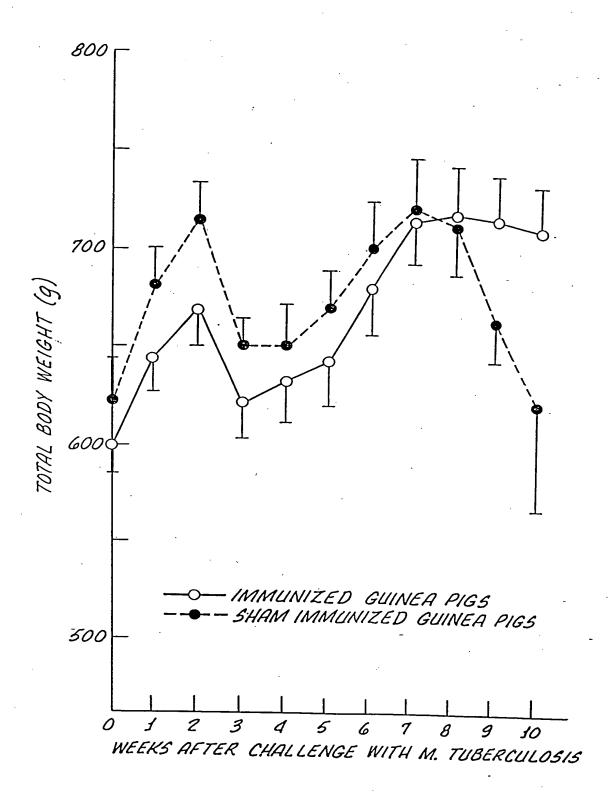
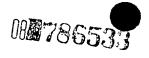


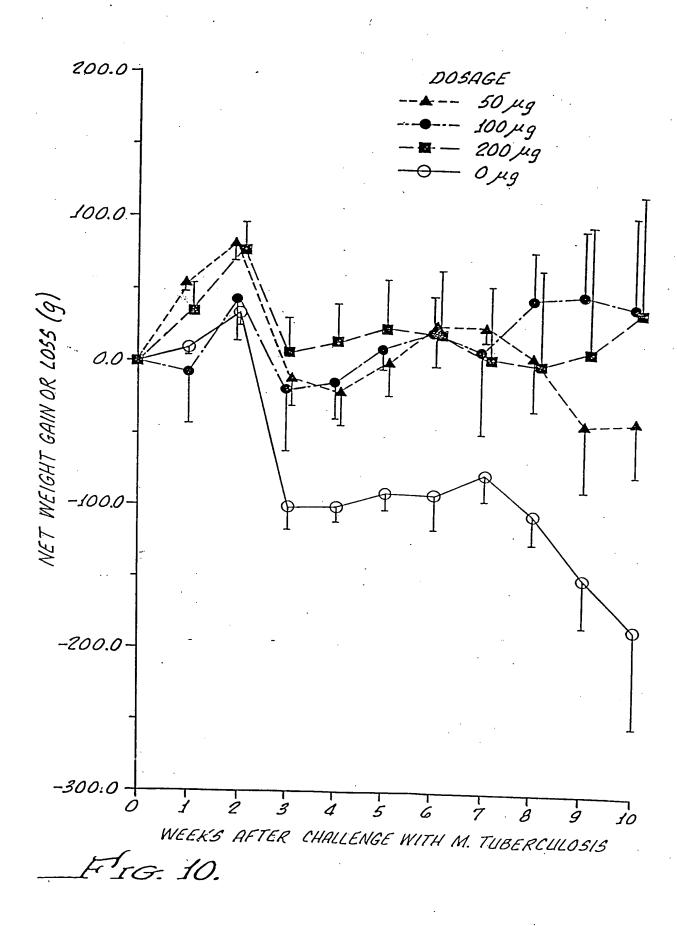
FIG. 7.

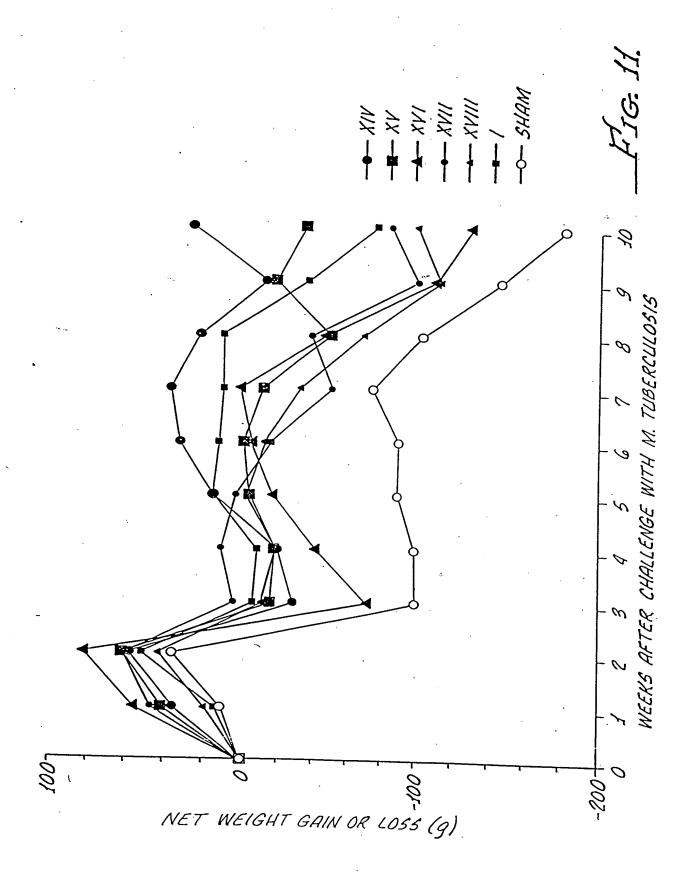
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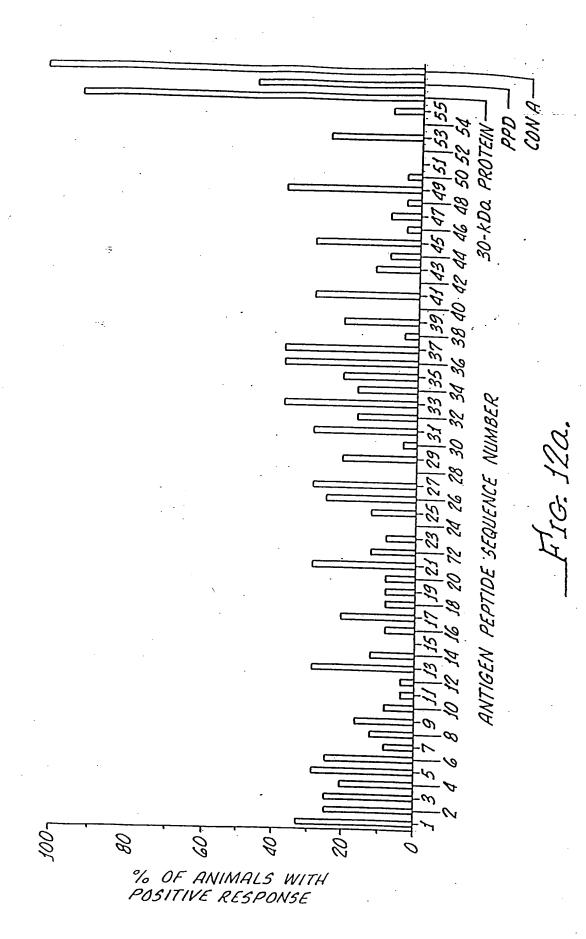


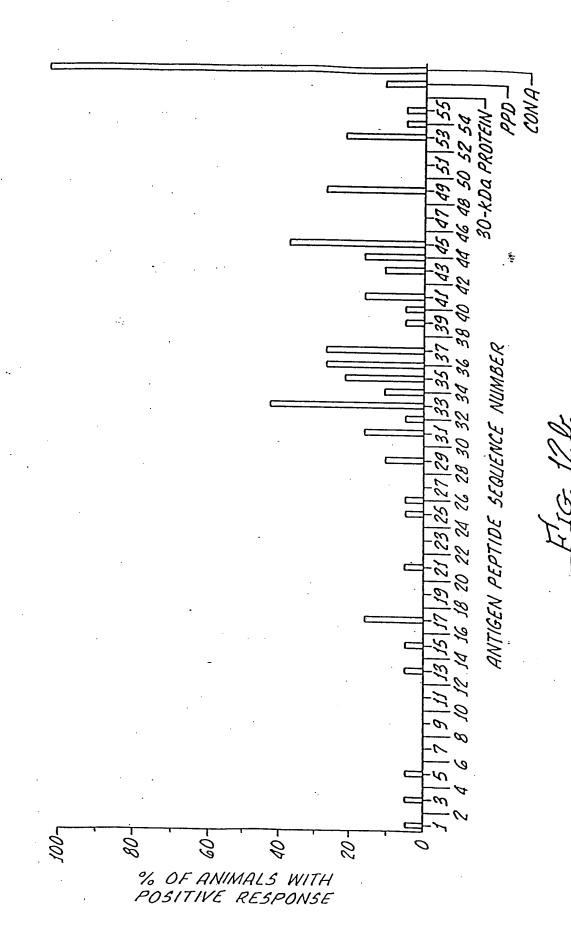
_FIG. 9.





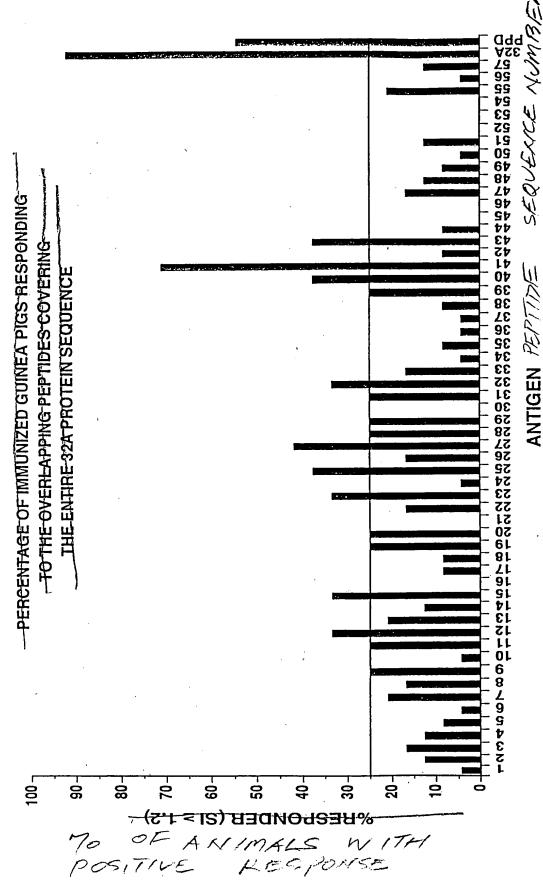






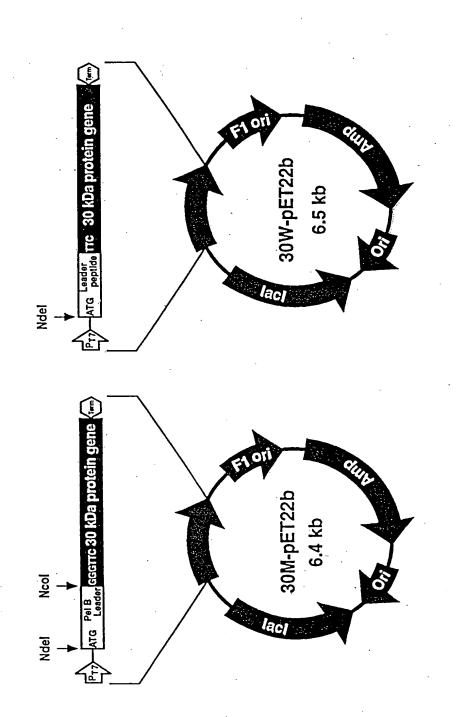
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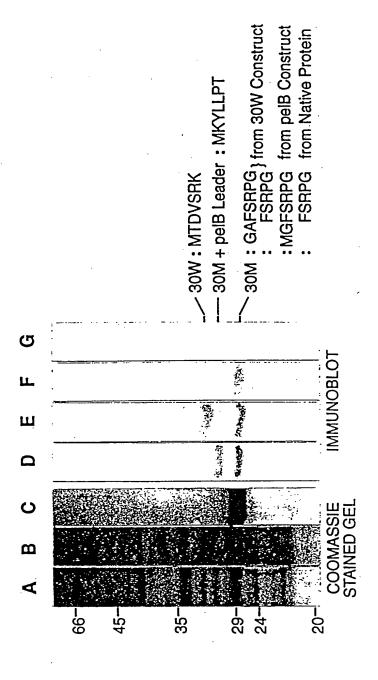


K19 13

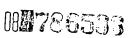




17 14



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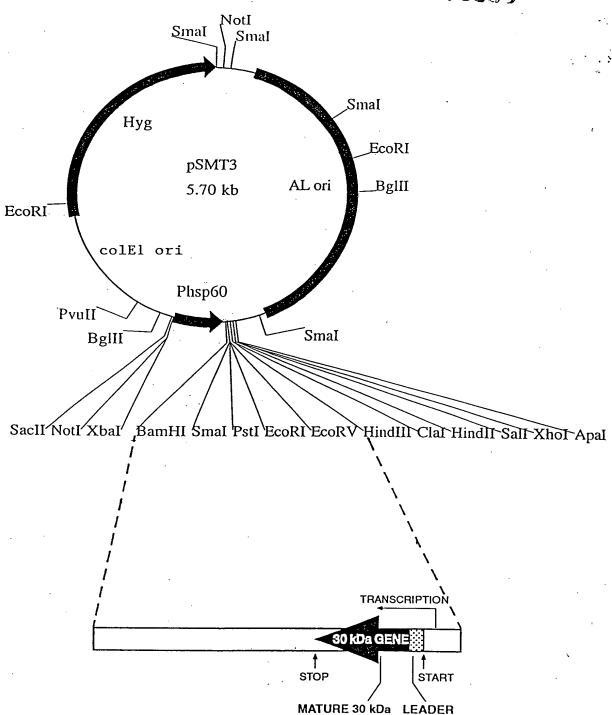
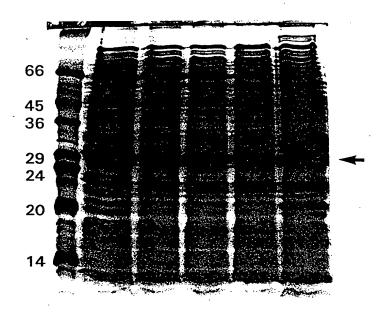


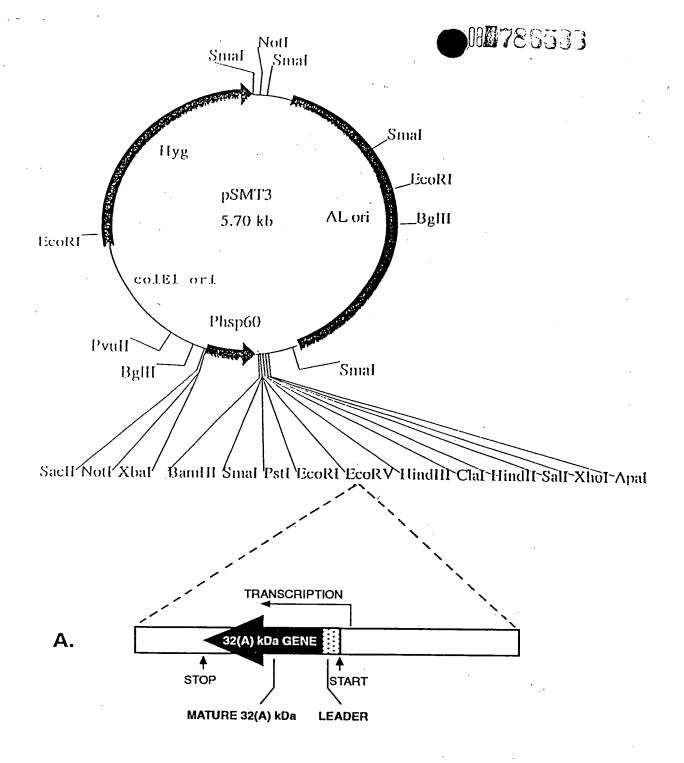
FIG. 16.

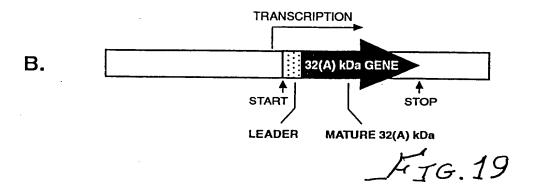


_FIG. 17.

106 80 49.5 32.5 27.5

F1G. 18.

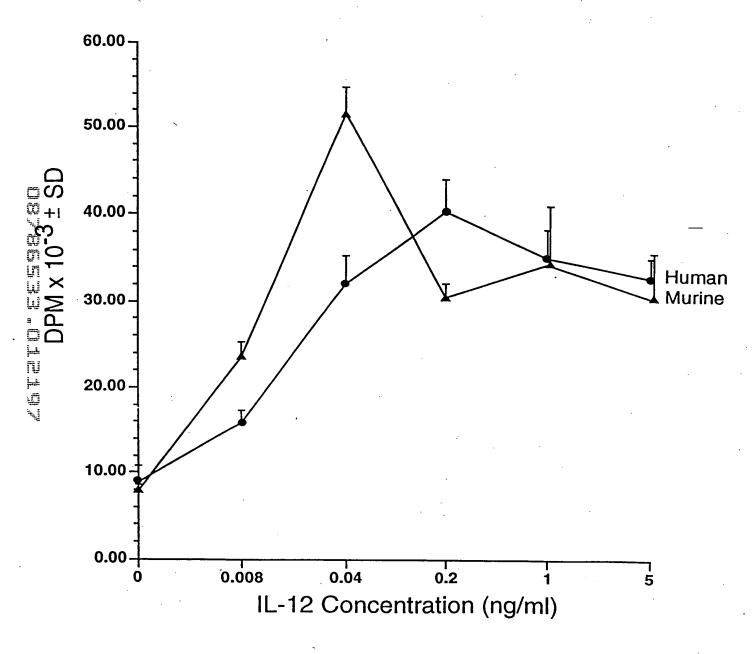




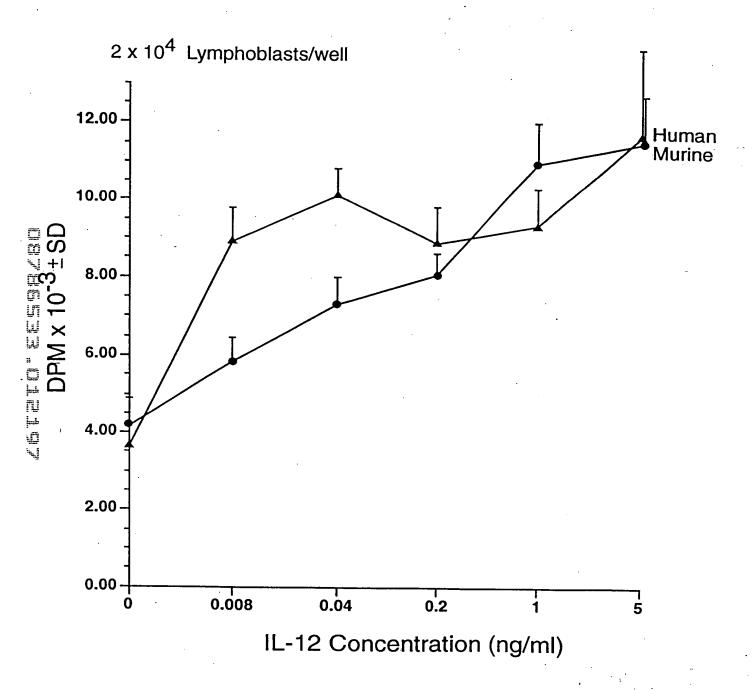
F1G. 20

IL-12 (Murine and Human)

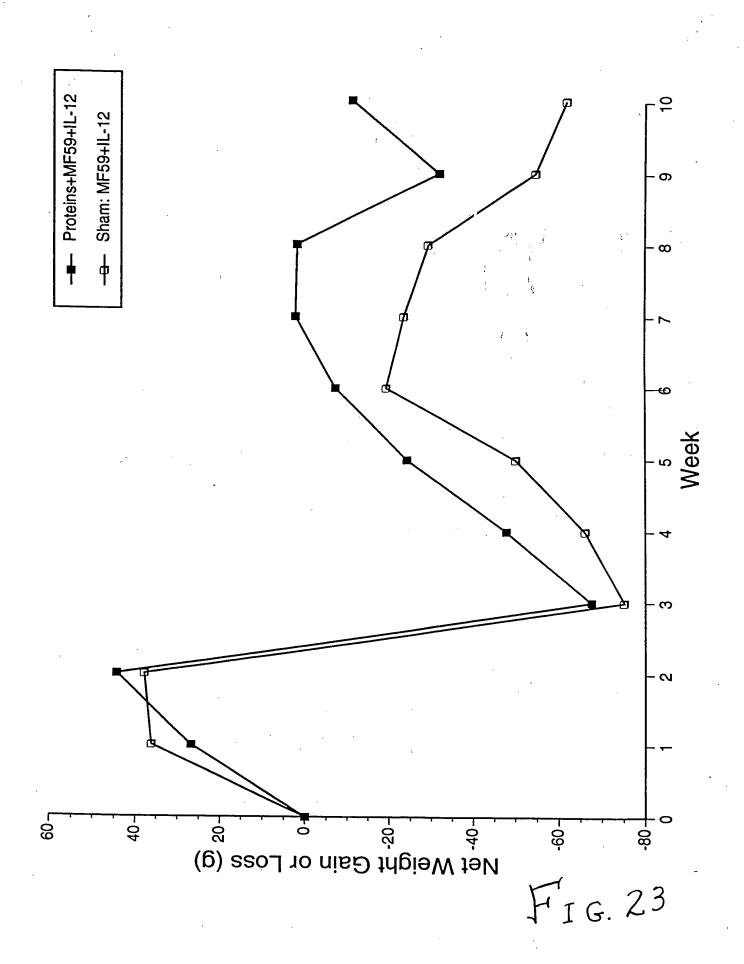
2 x 10⁴ Lymphoblasts/well

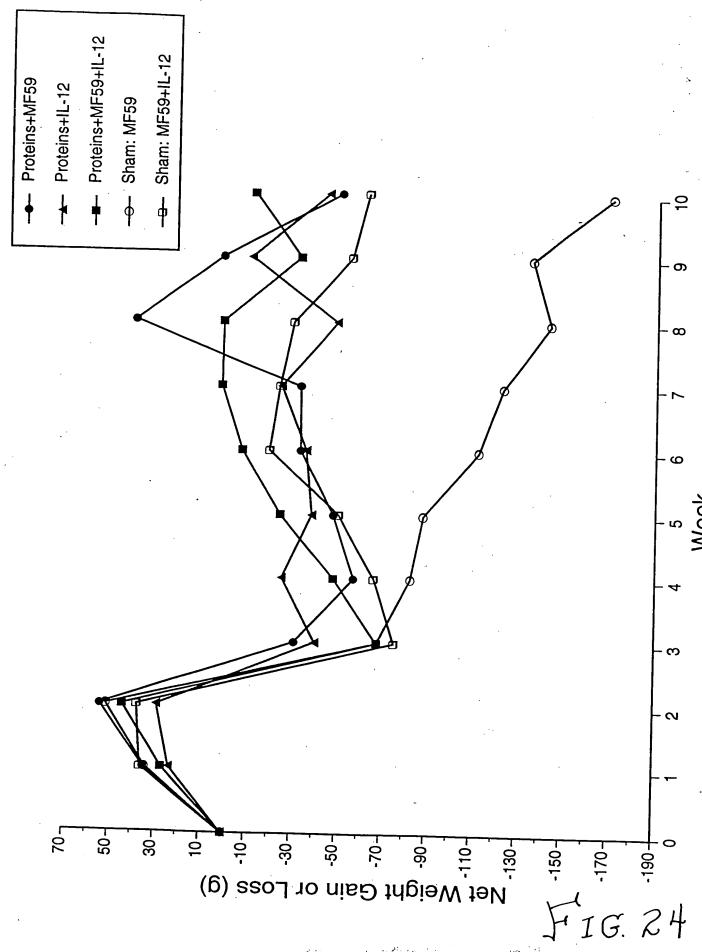


IL-12 (Murine and Human)



F1G. 22





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F.g. 25